SEQ ID NO.	BIALLELIC MARKER ID	ORIGINAL ALLELE	ALTERNATIVE ALLELE
1	20-828-311	С	T
1, 4	17-42-319	С	T
1, 2, 4	17-41-250	C	T
1	20-841-149	A	G
1	20-842-115	G	A
1	20-853-415	C	T

Figure 1

SEQ ID no.	Biallelic Marker ID	Original Allele	Alternative Allele	Position Range of Preferred Sequences
1	20-828-311	C	T	739-1739
1	17-42-319	С	Т	10946-12958; 13470- 13526; 13641-13752
1	17-41-250	C	Т	14271-17969
1	20-841-149	A	G	41718-42718
1	20-842-115	G	A	44942-45942
1	20-853-415	С	Т	76558-77558
2	17-41-250	С	T	1-1879
4	17-42-319	С	T	1-1498; 1613-1724
4	17-41-250	С	T	2243-3940; 3941-5381

Figure 2

A

SEQ ID NO.	POSITION OF CONFLICT	NUCLEOTIDE
1	13269 (SEQ ID No 1)	T (original)
4	1241 (SEQ ID No 4)	C (alternative)

B

SEQ ID NO.	POSITION OF CONFLICT	NUCLEOTIDE
1	13475 (SEQ ID No 1)	G (original)
4	1447 (SEQ ID No 4)	A (alternative)

Figures 3A, 3B

SEQ. ID. NO	POSITION RANGE O MICROSEQUENCING PRIMERS	COMPLEMENTARY POSITIN RANGE OF MICROSEQUENCING PRIMERS
1	1220-1238	1240-1258
1	12328-12346	12348-12366
1	15222-15240	15242-15260
1	42199-42217	42219-42237
1	45423-45441	45443-45461
1	77039-77057	77059-77077
4	300-318	320-338
4	3194-3212	3214-3232

Figure 4

SEQ. ID NO.	POSITION RANGE OF AMPLIFICATION PRIMERS	COMPLEMENTARY POSITION RANGE OF AMPLIFICATION PRIMERS
1	929-949	1357-1377
1	12029-12050	12581-12603
1	14992-15012	15460-15482
1	42070-42090	42572-42591
1	45328-45347	45863-45883
1	76644-76664	77166-77185
4	1-11022	553-11575
4	899-11920	1441-12461
4	1246-12267	1632-12651
4	2964-13984	3432-14454

Figure 5

SEQ. ID NO	POSITION RANGE OF PFOBES
1	1227-1251
1	12335-12359
1	15229-15253
1	42206-42230
1	45430-45454
1	77046-77070
4	307-331
4	3201-3225

Figure 6

Alignment of ApoA IV-related cDNA with Human and Swine cDNA's

ApoA IV related Human ApoA IV Swine ApoA IV	AGACCTGAGCAGAGCAGATAATCGCAAGCATGGCTGCCGTGCTCACCTGGGCT-CTTCTTTCAGCGTTTTTCGGC 79 AGTTCCCACTGCAGCGCAGGTG-AGCTCTCCTGAGGACCTCTCTGTCAGCTCCCCTGATTGTAGGGAGG 68
ApoA IV related Human ApoA IV Swine ApoA IV	CACCCAGGCACGGAAGGCTTCTGGGACTACTTCAGCCAGACCAGCGGGGACAAAGGCAGGGTGGAGCAGATCCATCAGC 159 CATCCAG-TGTGGCAAGAAACTCCTCCAGCCCAGCAAGCAGCT-CAGGATGTTCCTGAACCCAG-TGGAGTAAGAGAGACTTCCAGCCCAGCGGGAGCT-CAGGATGTTCCTGA93
ApoA IV related Human ApoA IV Swine ApoA IV	AGAAGATGGCTGGCGAGCCCGGAGCCCTGAAAGACAGCCTTGAGGAAGACCTCAACAATATGAACAAGTTCCTGGAAAAA 239 AGGGCGTGGTCC-TGACCCTGGCCCTGGTGGGTGTCGCCGGAGCCAGGGC-TGAGGTCAGTGCTGACCAGG 193 AGGCTGTGGTCC-TGAGCCTGGCCCTGGTGGCCGTCACCGGTGCCGGGCTGAGGTCAATGCCGACCAAG 162
ApoA IV related Human ApoA IV Swine ApoA IV	CTGAGGCQTCTGAGTGGGAGCQAGCTCCTCGGCTCCCACAGGACCCGGTGGGCATGCGGCAGCTGCAGGAGGAGTF] 319 TGGCCACAGTGAT-GTGGGACTACTTCAGCCAG-CTGAGCAGCAATGCCAAGGAGGCGGT-GGAACATCT 260 TGGCTACTGTGAT-GTGGGACTACTTCAGCCAG-CTGGGCAGCAATGCCAAGAAGGCTGT-GGAACATCT 229
ApoA IV related Human ApoA IV Swine ApoA IV	GGAGGAGGTGAAGGCTCGCGTCCAGCCCTACATGGCAGAGGCGGTACGAGCTGGTGGGGTGGAATTTGGAGGGCTTGCGGC 399 CCAG-AAATCTGAACTCACCCAGCAACTCAATGGCTCTTCCAGGACAAACTTGGAG
ApoA IV related Human ApoA IV Swine ApoA IV	AGCAACTGAAGCCCTACACGATGGATCTGATGGAGCAGGAGCTGCCCTTGCCGCGTGCAGGAGCTGCAGGAGCTTGCGCGTG 479AAGTGAACACTTACGCAGGAGCTGCAGAAGAAGCTGGTGCCCTTTGCCACCGAGCTGCATGAACGCCTG 387AAGTGAACACCTACACGGAGGACCTGCAGAAGAAGCTGGTGCCCTTTGCCACGGAGCTGCATGAACGCCTG 356
ApoA IV related Human ApoA IV Swine ApoA IV	GTGGGGGAAGAGAGAGAGGCCGAGTTGCTGGGGGGGGGG
ApoA IV related Human ApoA IV Swine ApoA IV	GETGCACCACACCCCCATACCCCCATACCCCGACACCCCGGCACACCCCGCACACCCCCATGCCCCACCGCCACCGCCACCGCCACCGCCCACCGCCACCGCCACCCCCACACCCCCACACCCCCACACCCCCACACCCCC
ApoA IV related Human ApoA IV Swine ApoA IV	CAGGAGTEGAACGGAAGTGTGCGGAAAGGCCCCAGGCAAGAGCCGGGGGGCGCCTCAGTGGCTGCGTGCAGAGTGGTGTGTGCGGGGGACAGCTGGGGACAGCTGGGACAGCTGGAGAGAGCAGCTGGAGAGAGCAGCTGGAGAGAGA
ApoA IV related Human ApoA IV Swine ApoA IV	GRAGOTOACOCTCAAGGOCAAGGOCTGCACGOAGGOAFICGAGCAGAACCTGGACCAGOTGGGGAAAGAGCTGAGCAGAG ACCCCTACGCACAGCGCATGGAGAGAGTGC-TGCGGGAGAACGCCGACAGCCTGCAGGCCTGAGGGGAGAG AAGCCCTACGCAGAGCGCATGGAGTCCGTGC-TACGGCAGAACATCCGCAACCTGGAGGCCTGCGTGGCA
ApoA IV related Human ApoA IV Swine ApoA IV	CONTI-TECRAGORACTICAGA AGGOCICAGORACTICA AGGOCICAGA AGGOCICAGA AGGOCICAGOCICAGA AGGOCICAGO
ApoA IV related Human ApoA IV Swine ApoA IV	GGTTTTCCGCAGGACACCTACCTGCAGATAGCTGCCTTCACTGGCCCATCGACCAGAGACTGAGGAGGTGCAGGAGAGG -GCCTTACGCCCTACGCTGAGGAATTCAAAGTCAAGATTGACCAGAGGGGGGGGGG
ApoA IV related Human ApoA IV Swine ApoA IV	AGCTTGCGCCCCTATGCTCAGGACACGCCGGGAGAAGCTCAACAAACAGACAGTGCCAAGCTTCTGAGCAAG 1037 AGCCTG-GCTCCCTATGCTCAGGACACGCAGGAGAAGCTCAACCACCAGCTTGAGGGCCTGA 838 AGCCTG-GCCCCCTATGCGCAGGACGTCCAGGAGAAGCTCAACCACCAGCTCGAGGGCCTGG
ApoA IV related Human ApoA IV Swine ApoA IV	CTGCAGGGCGTTTGGGATGACCTGTGGGAAGAATGACTCACAGGCTTCATGACCAGGGCGAGAGCCATGTGGGGGACCC 1117CCTTCCAGATGAAGAAGAACGCCGAGGAGCTCAAGGCCAGGATCTCGGCCAG-TG
ApoA IV related Human ApoA IV Swine ApoA IV	digagganciadececcaegocolatrocolagocicottetoeggagecttegetotaegocictaegategricaend 1197 cogaggagetegeegeagaegetegeeccottegecegaggaeetegetege-la 945 cogaggagetege
ApoA IV related Human ApoA IV Swine ApoA IV	CITCHANGTOCCTOTTOCCTOCAGAGGTCCTGTGCAGGACAGGGACACCAAAGGGGCTGCTGTGTCCTGC 1277 CTGAGGGCAACACCGAG-GGGCTGCAGAAGTCACTG-GCAGGAGGCTGGGGCACCTGGACCAGCAGGTGGAGG 1018 CTGAAGGGCAACACCGAG-GGGCTGCAGAAGTCGCTG-CTGGAGCTGAGAAGCCACCTGGACCAGCAGGTGGAGG 987
ApoA IV related Human ApoA IV Swine ApoA IV	ATATCCAGCCTCCCGGACTCCCCAATCTCACAACCCAGCCTTCCCAACCCAGCCTCCCAGTCCTCATT 1357 AGTTCCGACCCTCAACCCCTACGGGGAACCTTCAACAAACCCCTGGTGCAGCAGG 1047

Figure 7 A

Alignment of ApoA IV-related cDNA with Human and Swine cDNA's

ApoA IV related Human ApoA IV	TGGGPATGCTCATGAGTTACTCCATTGAAGGGTGAGGGAGTAGGGAGGG	1134
Swine ApoA IV	TGGAGGATCTCAGGC	
ApoA IV related	AAGGTGTTTTGCCGTGATGCTGGAAGGTCACTGCCACTACATCCTGGAGTTTGGCTCAGTGATGACTTTCTGGCTGCCTGGTGGGAGAAGGACAAGGTCA	
Human ApoA IV Swine ApoA IV	-AGCTTCCTGGAGAAGGATCTGAGGGACAAGGTCAACACCTTCTTCAG-CACCTTCAA	
ApoA IV related	CCACTGCTACAGCTGGTCCACAGAGAGCTAGTGTGTCCCCAGGGTTGCCATGCAGCTATCAGGGGAATAGAAGGGA AAGAGAGCCA-GGACAAGACT-CTCCCTCCCTCAGCTGGAGCAACAGCAGGAAGAGCAACAGCATCAG	1597
Human ApoA IV Swine ApoA IV		1199
ApoA IV related	GAAAGAATATCATGAGAAAAAAATATCATGAGAAAAATATCCCTGCTCTGATGATGAGAAAAAGAGAAAAAGAGAAAAAGAGAAAAAGAGAAAAA	
Human ApoA IV Swine ApoA IV	CAGGAGAAAGGGCAGGCCCCTTTGGAGGGCTGAGCTGCCCCTGGTGCTCCCACCCCAC	
ApoA IV related	TGGGTTGTGATAGGAGAGGGCAGAGCCCATGTTTCCTGAGATAGGTTTACACCTAAATAAGGGACTGAACCCTCCCAACT	1757
Human ApoA IV		1376
Swine ApoA IV	ACGTCTGT-CTGTCCCAA	1304
ApoA IV related	eredekedridirinaakcciicideedaecirikcrerereciorecceirioreckedocicorreceiriodeakeriig	
Human ApoA IV	adaagttc-iggtatgaactidaggacacatgtccagtgggaggtgagaccacctctdataticcaata agdagttc-ttgtacaaaddagggatacatgtccagtggaccgtgadactacctctddatactcaata	
Swine ApoA IV		1,7,2
ApoA IV related		1879
Human ApoA IV		1466
Swine ApoA IV	AAGCTGCTGAGAAACT	1388

Alignment of ApoA IV-related protein with Human and Swine ApoA IV

NKFLEKI 74 GDLQKKU 80 EDLQKKI 80	GEOTRVV 154 RRQLTPY 160 ORQLKPY 160	ISRCVQV 230 LNHQLEG 240 LNHQLEG 240	FIRMIDG 305 FNKALVG 320 FNKALVG 320	366 LES 396
MASMAAVIJNMALALLSAFSATGARKGFWDYFSGTGGD-RGRVBQIHQQKMAREPATI-KDSLTEQDLNNMNKFLEKI	RPLSGSERAPRIPQPPVGMRRQLQEELEBVKARIQPYMABAHELVGWNIEGIRQQHKPYTMDTMEQVALRVGELGEQIRVV	GBDTKAQILIGGVDEAWALIQGIQSRVVHHTGRFKBLFHPYABSILVSGIJGRHVQETHRSVAPHAPASPARIJSRCVQV	SRKLTLKARALHARTQONLGOLREBLISRAFAGTGTEBGAGPDPQMLSBEVRGRLQAFRQDTYLQIAAFTRAIDG	ЕТВЕVOGOЦАВРРРЕНSAFAPEFQTDSGKVLSKLQARLDDLWEDITHSU
MFIKAVVITLALVAVAGARAEVSADQVATVMWDYFSQLGNNAKBAVEHLQKSEITQQINALFQDKLGEVNTYAGDLQKKI	VPFATELHERIAKDSEKLKEETGKELEELRARLLPHANEVSQKIGDNLRELQQRIEPYADQLRTQVNTGAEQIRRQLTPY	AQRMBRVLRBNADSILQASIRPHADELKAKIDQNVEELKGRLTPYADBFKVKIDQTVBELRRSLAPYAQITQEKLNHQLEG	TFQMKKNABEBLKARISASABBLRGRLARIABDVRGNIRGNTEGLOKSIJABIGGHLDQOVBEFRRRVEPYGBNFNKALVQ	
MFIKAVVISLALVAVTGARAEVNADQVATVMWDYFSQIGSNAKKAVEHLQKSEITQQINTLFQDKLGEVNTYTEDIQKKI	VPFATELHERITKDSEKLKEETRRELEELRARLLPHATEVSQKIGDNVRELQQRIGHFTGGLRTQVNTQVQQLQRQIKPY	AERMBSVLRQNIRNLBASVAPYADBFKAKIDQNVEELKGSLTPYABELKAKIDGNVBELRRSLAPYAQDVQEKLNHQLEG	AFQMKKQABBLKAKISANADBLRGKLYBVABNYHGHLKGNTEGLQKSYLBIRSHLDQQVEBFRLKVEPYGBTFNKALVQ	
MASMAR MFLKAV	RPLSGS VPFATE VPFATE	GEDTKA AQRMER AERMES	LSRKLT LTFOMK LAFOMK	ETBEVQ QMEQLR QVEDLR
ApoA IV related	ApoA IV related	ApoA IV related	ApoA IV related	ApoA IV related
Human ApoA IV	Human ApoA IV	Human ApoA IV	Human ApoA IV	Human ApoA IV
Swine ApoA IV	Swine ApoA IV	Swine ApoA IV	Swine ApoA IV	Swine ApoA IV

Figure 8

Alignment of ApoA IV-related cDNA with Rat RAP3 cDNAs

ApoA IV related	AGACGTGAGCAGAGAGATAATGGCAAGCATGGCTGCCGTGCTCACCTGGGCTCTGGCTCTTCTTTCAGCGTTTTCGGCC	80
Rat RAP3 A	GCATCGTGGAAAGCATGGCTGCCGTCATCACCTGGGCACTCGCCCTCCTCTCAGTGTTTGCAACT	65
Rat RAP3 B	GCATCGTGGAAAGCATGGCTGCCGTCATCACCTGGGCACTCGCCCTCTCTCAGTGTTTGCAACT	65
ApoA IV related	ACCCAGGCACGGAAAGGCTTCTGGGACTACTTCAGCCAGACCAGGCGGGACAAAGGCAGGTGGAGCAGATCCATCAGCA	160
Rat RAP3 A	GTACAGGCGAGGAAGAGCTTCTGGGAGTACTTCGGCCAGAACAGCCAGGCAAAGGCATGATGGGCCAGCAGCA	139
Rat RAP3 B	GTACAGGCGAGGAAGAGCTTCTGGGAGTACTTCGGCCAGAACAGCCAGGGCAAAGGCATGATGGGCCAGCAGCA	139
ApoA IV related	GAAGATGGCTTGGGGAGCCCGCGAGCCTGAAAGACAGCCTTGAGCAAGACCTGAACAATATGAACAAGTTCCTTGGAAAAAGC	240
Rat RAP3 A	GAAGCTGGCACAGGAGAGCCTGAAAGGTAGCTTGGAGCAAGACCTCTACAATATGAACAATTTCCTAGAAAAAGC	213
Rat RAP3 B	GAAGCTGGCACAGGAGAGCCTGAAAGGTAGCTTGGAGCAAGACCTCTACAATATGAACAATTTCCTAGAAAAGC	213
ApoA IV related	TGAGGCCTCTGAGTGGGAGCGAGGCTCCTCGGCTCCCACAGGACCGGTGGGCATGCGGCAGCTGCAGGAG	. 274
Rat RAP3 A	TGGGACCCTTGAGAGAGCCTGGGAAGGAGCCTCCTCGGCTGGCACAGGATCCAGAAGGCATTCGGAAGCAGTTGCAGCAA	202
Rat RAP3 B	TGGGACCCTTGAGAGAGCCTGGGAAGGAGCCTCCTCGGCTGGCACAGGATCCAGAAGGCATTCGGAAGCAGTTGCAGCAA	293
ApoA IV related	GACTTGGAGGAGGTGAAGGCTCCAGCCCTACATGGCAGAGGCGCAGGAGCTGGGGTGGGCTTGGAATTTGGAGGGCTT	
Rat RAP3 A	GAGCTGGAGGAAGTGAGCACACGCCTGGAGCCCTACATGGCTGCAAAGCACCAGCAGGTCGGCTGGAACCTGGAGGGCTT	394
Rat RAP3 B	GAGCTGGAGGAAGTGAGCACACGCTGGAGGCCTACATGGCTGCAAAGCACCAGCAGGTCGGCTGGAACCTGGAGGGCTT	373 373
ApoA IV related	GOGGCAGCAAGTGAAGCCCTACACGATGGATCTGATGGAGCAGGTGGCCCTGGGGGTGCAGGAGCTGCAGGAGCAGTTGG	474
Rat RAP3 A	GAGGCAGCAGTTGAAACCCTACACGGTCGAGCTGATGGAGCAGGTAGGCCTGAGCGTGCAGGATCTGCAAGAACAGCTGC	453
Rat RAP3 B	GAGGCAGCAGTTGAAACCCTACACGGTCGAGCTGATGGAGCAGGTAGGCCTGAGCGTGCAGGATCTGCAAGAACAGCTGC	453
ApoA IV related	GdGTGGTGGGGAAGACACCAAGGCCCAGTTGCTGGGGGGGGG	554
Rat RAP3 A	GCATGGTGGGAAAAGGCACCAAGGCCCAGCTCCTGGGGGGCGTGGATGAGGCGATGAGCCTGCTGCAGGATATGCAAAGT	533
Rat RAP3 B	GCATGGTGGGAAAAGGCACCAAGGCCCAGCTCCTGGGGGGGG	533
ApoA IV related	cdcTcGTcCAccAcAdcGcCcCTTCAAAGACCTCTTCCACCCATACGCCGAGAGCCTGGTGAGCGCATTGGGGCCCA	634
Rat RAP3 A	CGAGTGCTGCACCATACGGACCGAGTCAAAGAACTCTTCCACCCTTATGCAGAACGCTTGGTGACTGGAATTGGGCACCA	613
Rat RAP3 B	CGAGTGCTGCACCATACGGACCGAGTCAAAGAACTCTTCCACCCTTATGCAGAACGCTTGGTGACTGGAATTGGGCACCA	613
ApoA IV related	detecagaectecaccedaeteridectcoecacedecccaecccecedecctcaetecatecaetecae	714
Rat RAP3 A	TGTGCAGGAGCTGCACCGGAGTGTTGCTCCTCACGCAGTTGCCAGCCCCGCGAGACTCAGTCGCTGCAGGACCCCGT	693
Rat RAP3 B	TGTGCAGGAGCTGCACCGGAGTGTTGCTCCTCACGCAGTTGCCAGCCCCGCGAGACTCAGTCGCTGCAGGACCCTGT	693
A A . 137 1.4 3		
ApoA IV related	CCCCCCAAGCTCACCCTCAAGCCCAAGCAACCTCCACCAAGCAAACCTCCACCAAGCAACCTCCCCCAAACCAACC	794
Rat RAP3 A	CCCACAAACTCACACGTAAGGCGAAGGACTTGCACACCAGCATCCAACGCAACCTGGATCAGCTGCGAGATGAGCTCAGT	773
Rat RAP3 B	CCCACAAACTCACACGTAAGGCGAAGGACTTGCACACCAGCATCCAACGCAACCTGGATCAGCTGCGAGATGAGCTCAGT	773
ApoA IV related		868
Rat RAP3 A	ACCTTCATCCGTGTCAGCACAGACGGGGCAGACAACAGAGACTCCCTGGACCCTCAAGCTCTCTCT	853
Rat RAP3 B	ACCTTCATCCGTGTCAGCACAGACGGGGCAGACAACAGAGACTCCCTGGACCCTCAAGCTCTCTGGACGAGGTCCGCCA	853
ApoA IV related	GCGACTICAGGCTTICCGCCAGGACACCTACCTGCAGATAGCTGCCTTCACTCGCGCCCATCGACCAGGAGACTGAGGAGG	948
Rat RAP3 A	GAGACTCCAGGCTTTTCGACATGACACCTACCTGCAGATCGCTGCATTCACTCAGGCCATTGACCAGGAGACCGAGGAAA	933
Rat RAP3 B	GAGACTCCAGGCTTTTCGACATGACACCTACCTGCAGATCGCTGCATTCACTCAGGCCATTGACCAGGAGACCGAGGAAA	933
ApoA IV related	TCCAGCAGCAGCTGGGGCCACGTGGACCACGCCCACAGTGCCTTCGCCCCAGAGTTTCAACAAACA	1028
Rat RAP3 A	TCCAGCACCAGCTGGCACCACCCCCGCCTAGCCACAGCGCCTTCGCTCCAGAGTTGGGACACTCAGACAGTAATAAGGCC	1013
Rat RAP3 B	TCCAGCACCAGCTGGCACCACCCCCGCCTAGCCACAGCGCCTTCGCTCCAGAGTTGGGACACTCAGACAGTAATAAGGCC	1013
poA IV related	CTGAGCAAGCTGCAGGCCCTTCTGGATGACCTGTGGGAAGACATCACTCAC	1100
at RAP3 A	CTGAGCAGACTGCAGAGCCGGCTGGACGACCTCTGGGAAGATATTGCCTATGGCCTTCATGACCAGGGCCATAGTCA	1090
at RAP3 B	CTGAGCAGACTGCAGAGCCGGCTGGACGACCTCTGGGAAGATATTGCCTATGGCCTTCATGACCAGGGCCATAGTCA	1090
poA IV related	GGGGGACCCCTGAGGATGTACCTGGCCAGGCCGATTGC-CAGCTCCTTGTCTGGGGAGCCTTGGCTCTGAGCCTCTAGCA	1107
at RAP3 A	-GAATAACCCTGAGGGTCACTCAGGTTAACTCTGCAGCTCGTTGTCTGGACCCTGAGCCTCTAGCA	1155
at RAP3 B	-GAATAACCCTGAGGGTCACTCAGGTTAACTCTGCAGCTCGTTGTCTGGACCCTGAGCCTTCAGCA	1155

Figure 9 A

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Alignment of ApoA IV-related cDNA with Rat RAP3 cDNAs

ApoA IV related Rat RAP3 A	TGGTTCAGTCCTTGAAAGTGGCCIGTTGGGTGGAGGGTGGAGGGTCCTGTGCAGGACAGG-GAGGCCACCAAAGGGCTG TGGCCTAATAGGCAGAGGGTGGAGGGTCCTGCATACTATTGGGGAGGCCACCAAAGGTGCTG	1217
Rat RAP3 B	TGGCCTAATAGGCAGAGGGTGGAGGGTCCTGCATACTATTGGCGAGGCCACCAAAGGTGCTG	1217
ApoA IV related Rat RAP3 A Rat RAP3 B	CTGTCTCGTGCATATCCAGCCTCCTGGGACTCCCGAATCTGGATGCATTACATTCACCAGGCTTTGCAAAGCCAGGCTCCCTG-CCCCAACCTGTCTGGCCACCCCACC	1346 1295 1285
ApoA IV related Rat RAP3 A Rat RAP3 B	CAGTGCTCATTTGGGAATGCTCATGCATTCAAGGGTGAGGGAGTAGGGAGGAGAGGCACCATGCATG	1426 1361 1285
ApoA IV related Rat RAP3 A Rat RAP3 B	TGATTATCTGCAAGCCTGTTTGCCGTGATGCTGGAAGCCTGTGCCACTACATCCTGGAGTTTGGCTCTAGTCACTTCT AGACTGACTGCAAGCCAGTACTTGAC-CGTTGCTAGAAACCTGTGTCACTACAACCTGGAGCCCGGCTCCTATTACTTCA	1504 1440 1285
ApoA IV related Rat RAP3 A Rat RAP3 B	GGTIGCCTGGTGGCAGACTGGTACAGCTGGTCACAGAGAGAGGAGCACTTGTCTCCCCAGGGCTGCCATGGCAGCTATICAGG TGCCTGATGGTGGCTGTTATAGTCGGTGTACAGAGGGGAAGTCCTGTCTCCCCCAGGGTTGTCATGACAGGCTTTGTT 	1584 1517 1285
ApoA IV related Rat RAP3 A Rat RAP3 B	GGAATAGAAGGGAGAAAGAGAATATCATGGGGAGAACATGTGATGGTGTGTGAATATCCCTGCTGGGTCTGATG GGAAGAGGAGCAGGAGAACATGACACGTATGATGGAGTGTGTACATCCCTGCGGGTCCTGCTGGGGGAATCAGTGATG	1658 1597 1285
ApoA IV related Rat RAP3 A Rat RAP3 B	ctggtgggtaddaaacgtgtgggdtgtgataggagggggggggggggggtgggtgttcctdadata ggataaatgtgtggatccctgdagtggtcctgdtgggggatdagtgataggatag	1720 1677 1285
ApoA IV related Rat RAP3 A Rat RAP3 B		1799 1754 1285
ApoA IV related Rat RAP3 A Rat RAP3 B	TOTTCATCATCCACACTCCCGATTCTCGGGATCGAAGTGACTTTTGGTTGGAATGAAATAGACGCTCATCAT	1878 1832 1285
ApoA IV related Rat RAP3 A Rat RAP3 B	GGAAAAAAAAAAAA 	1879 1850 1300

owerset etest

Alignment of ApoA IV-related protein to Rat RAP3 proteins

PLLES 78	VGEDT 158	KLTLK 238	20001A 315	366
PLREP 73	VGKGT 153	KLTRK 233	2HQLA 312	367
PLREP 73	VGKGT 153	KLTRK 233	2HQLA 312	367
MASMAAVITWALALISARISATQAR KGFWIDYFISQIKGRVEQIHQQKMAREPATLKIDS LEQDINNMNKFLEKLRPIL-S	GSBAPRUROPPUGMRROLOBBLEEVKARLIOPYMABAHBLIVGWNLEGLROOLKPYTMDLMEOVALBRVOBLOEGLRYVGBDT	KAQLLGGVDEAWALLQGLQSRVVHHTGREKELFHPYABISLVISGI GRHVQELHRSVAPHAPASPARLSRCVQVLSRKLTLK	AKALHARIOONIDOIRBEISRAR – – AGIGIBEGAGPD POMLSBEVRORIOAFRODTYLOIAAFTRAIDOETEEVOOLA	PPPPGHSAFAPEFQQTDSGKVLSKLOARLDDLWEDITHSLHDQGHSHLGDP
MAAVITWALALISVFATVQARKSFWEYFGQNSQGKGMMGQQQKLAQESLKGSLEQDLYNMNFLEKLGPLREP	GKEPPRLAODPEGIRKOLOOBLEEVSTRLEPYMAAKHOOVGWNLEGLROOLKPYTVELMEOVGLSVODLOEGLRWGKGT	KAQLLGGVDEAMSLLQDMQSRVLHHTDRVKELFHPYABRLVTGI GHHVQELHRSVAPHAVASPARLSRCVQTLSHKLTRK	AKDIHTSIORNIDOIRDEIS-TFIRVSTDGADNRDSIDPOALSDEVRORIOAFRHDTYLOIAAFTOAIDOETEEIOHOLA	PPPPSHSAFAPELGHSDSNKALSRLQSRLDDLWEDIAYGLHDQGHSQNNPEGHSG
MAAVITWALALISVFATVQARKSFWEYFGQNSQGKGMMGQQQKLAQESLKGSLEQDLYNMNFLEKLGPLREP	GKEPPRLAODPEGIRKOLOOSLEEVSTRLEPYMAAKHOOVGWNLEGLROOLKPYTVELMEOVGLSVODLOEGLRWYGKGT	KAQLLGGVDBAMSLLQDMQSRVLHHTDRVKELFHPYABRLVTGI GHHVQELHRSVAPHAVASPARLSRCVQTLSHKLTRK	AKDIHTSIORNIDOIRDEIS-TFIRVSTDGADNRDSIDPOALSDEVRORIOAFRHDIYLOIAAFTOAIDOETEEIOHOLA	PPPPSHSAFAPELGHSDSNKALSRLQSRLDDLWEDIAYGLHDQGHSQNNPEGHSG
Rat RAP3 A Rat RAP3 B	ApoA IV related Rat RAP3 A Rat RAP3 B	ApoA IV related Rat RAP3 A Rat RAP3 B	ApoA IV related Rat RAP3 A Rat RAP3 B	ApoA IV related Rat RAP3 A Rat RAP3 B

Figure 10

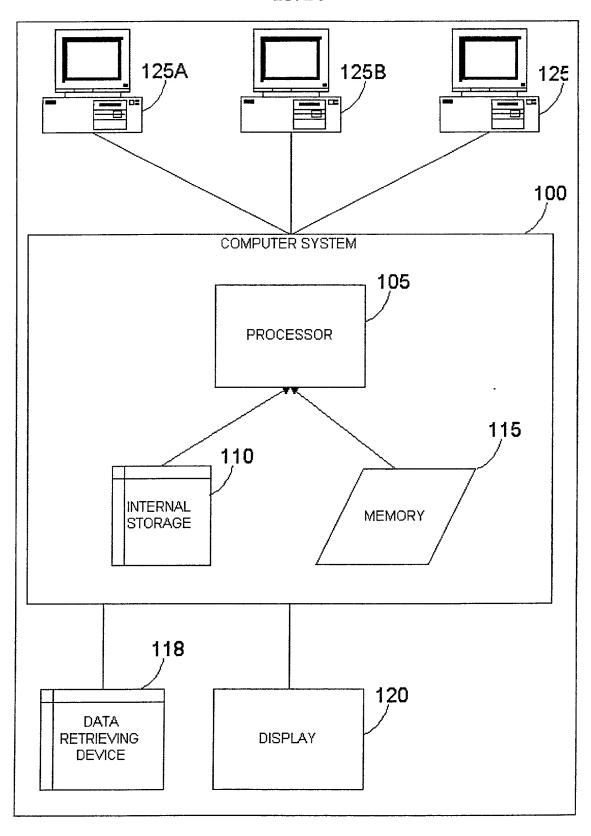


Figure 11

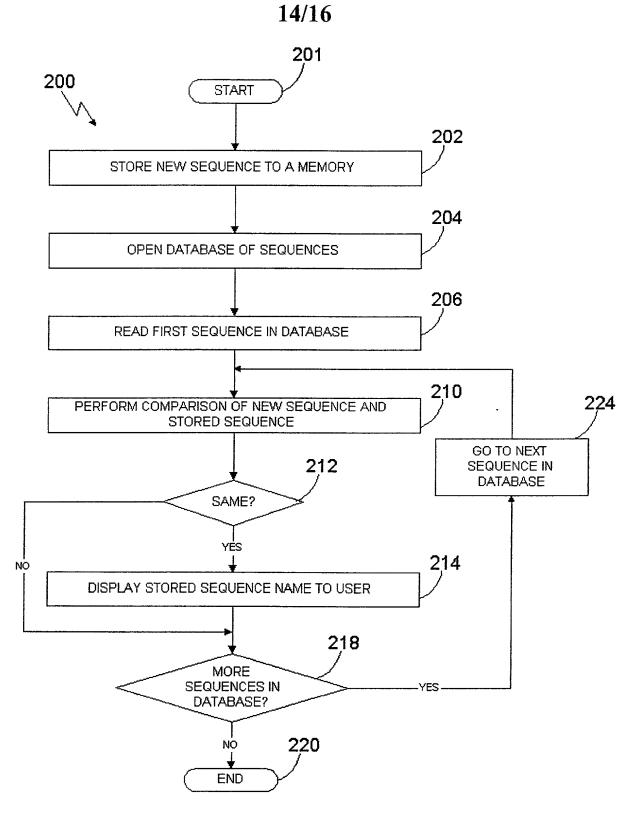


Figure 12

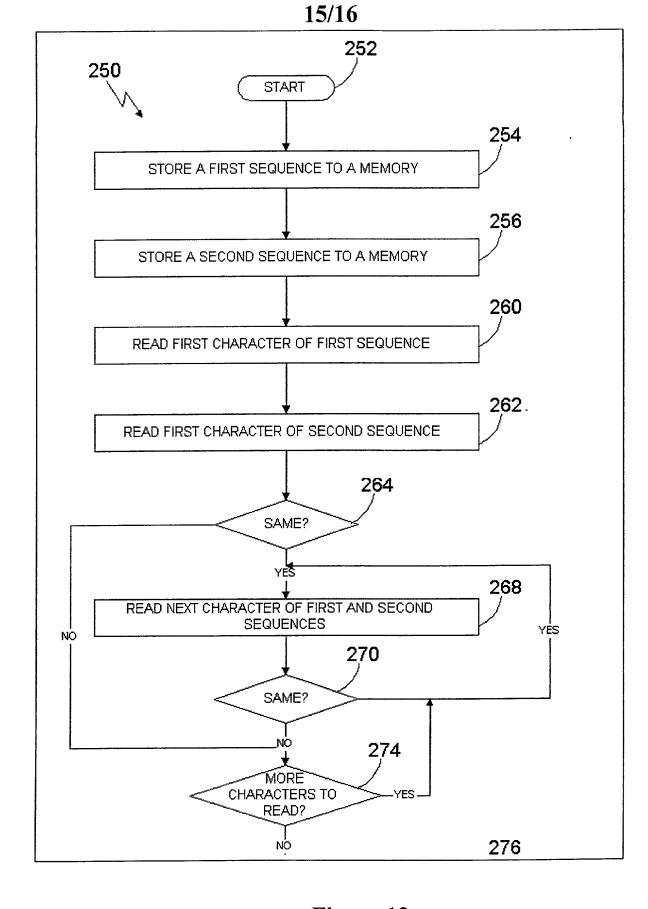
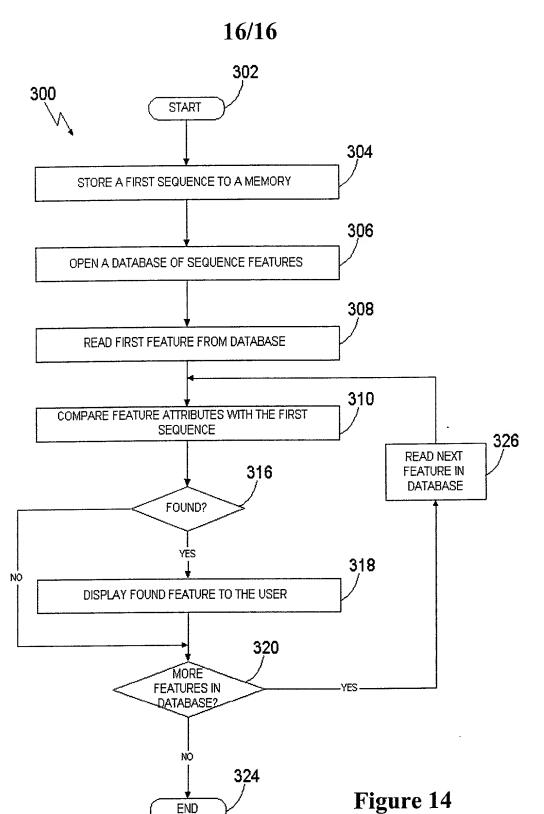


Figure 13



END